Figure 1

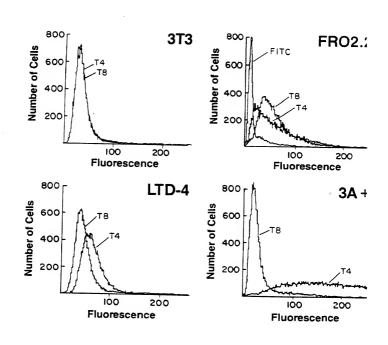
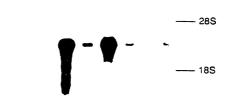


Figure 2

L cell LTD-4 tk7 HeLa IMR SK-7 OT-CLL Fro 2.2 T4 *T cells T4 *T cells





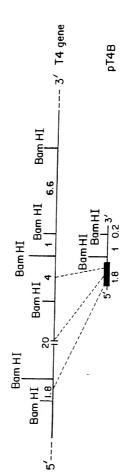


Figure 3B

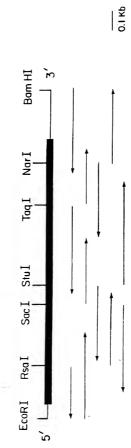
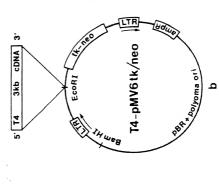
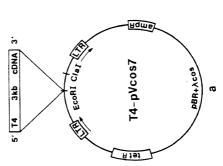


Figure 30





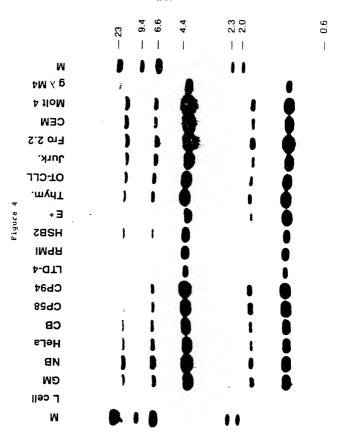


Figure 5

25.7 —

Figure 6A

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Arg me	468	A H	asn MC	G Pa	88	Ag th	val GTC	ug.
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Figure 6B

9	Leu the leu pro gln ala leu pro gln tyr ala gly ser gly asn leu thr leu ala leu gly alv ala lys thr gly lys leu his gln glu CCC ACC CTG CCC CAG CCC TGC CTC CAG TAC CCT CGC TCT CGC ACC CTT CGA CAG AAA CAG CGC AAA ACC CTC CAG CAG CAT CGA AAA CAG CAG AAA CAG CAG CAC CAG CGC CTT CGA CAG AAAA CAG CAG AAA CAG CAG CAC CAC	val and lea val val met arg ala thr gln leu gln lys aen leu thr cys glu val trp gly pro thr ser pro lys leu met leu ser gro Anc cor cor cor cor cor cor cor cor cor co	1188	1278	1368	HIO HIO BY AND HIS LYS AND HIS LYS BY DIS COST ONE COST TIT ONE AND NOT THE COST TIT ONE AND NOT	
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+230 Lib thr phe asp leu lys asm lys glu val ser val lys arg val thr glm asp pro lys leu glm met gly lys lys leu pro leu his ARC ACC TIT GAC CTUS ANG AAA GAA GTA GTA AAA CGTA AAC CTU AAG CTUC CTUG ANG GTC AAC AAA CTA CTA CTA CTA CTUC CTUG ANG GTC CTA CAAC CTA CTA	35	13 2	+320 +340 be by a sen lys glu ala lys val ser lys arg glu lys ala val trp val leu asn pro glu ala gly met trp gln cys leu rro AMA CTG GMG AMC AMG GGG GCG AMG GGC GGG GGG GGG GGG GGG GGG GGG GGG G	+350 +350 +370 +301 Let set asp set gly gln val let let glu set asn ile lys val let pro thr trp set thr pro val gln pro met ala let ile val CTG NOT GNC TOG GOA GNC GTG GNA TVC ANC ANT GTG GCC ANA TVC GTG GNC GNC GNC GNC GNC GNC GNC GNC GNC GN	+380 +380 +480 480 480 480 480 480 480 4	4 4 6	_
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9/30

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осм'ятактыстабая**тмозоскосс**тусмстватурм фатролест (тубатуюст учество поставляет в 1897)

BP T4

11/30

Figure 8

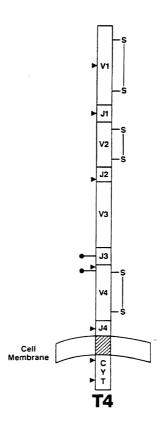


Figure 9A

12/30	
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	1
τ4 V-kappa Inv. T8 XT35 HPB-HLTΩ	T4 V-kappa Inv. T8 YT35 HPB-MLT α

Figure 9B

 $\begin{array}{c} 100 \\ \text{T4 J} \\ \text{Consensus } JB \\ \text{Consensus } JX \\ \text{Consensus } JK \\ \text{HPB-MIT } \alpha \text{ J} \\ \end{array} \begin{array}{c} 100 \\ \text{L V F G L } - \text{TA N S D T H} \\ \text{L Y F G E G T R L T V L } - \\ \text{W V F G G G T K V T V L G} \\ \text{W T F G G G T K L E I K R} \\ \text{HPB S G T R L S I R } - \\ \end{array}$

Figure 90

370 STPVOPMALIVLGGVAGLLLFIGLGIFFCVR STSAQNKMLSGVGGFVLGLLFLGLGLELFIYFR

> 4 HC cl.11β

Figure 10

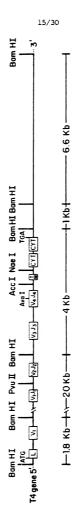
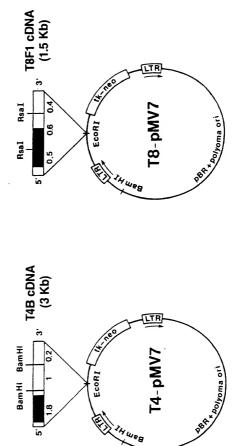


Figure 11A



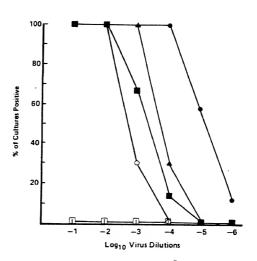
ECOR!

BamHi

Figure 11B

	MOUSE TRANSFORMANT	HUMAN TRANSFORMANIS HELA Carcinoma HSB2 T cells RPMI B cell
Colony Purification	<u>MOUSE</u> Stable T4*(or T8*)	Stable T4*(or T8*)
Rosetting	Purification	Purification
G418 Selection	G418 Selection	G418 Selection
√ -2	Infection Murine cells	Co-cultivation Human cells
10 µg T4-pMV7 Transfect (or T8-pMV7) DNA	14 (or 18 1)	T4 for T8 +) Mitomycin C W-AM

Figure 12



19/30

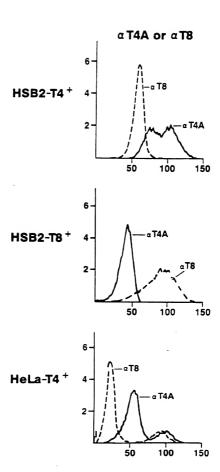
Figure 13A

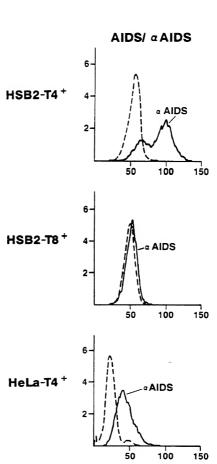


Figure 13B



21/30 Figure 14A





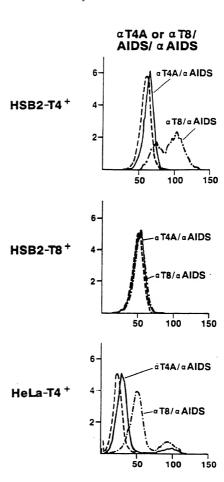


Figure 15A

RAJI U937 Jurkat Cortex

-28S



Figure 15B

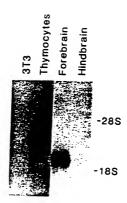
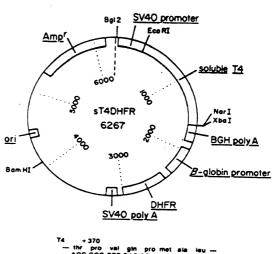
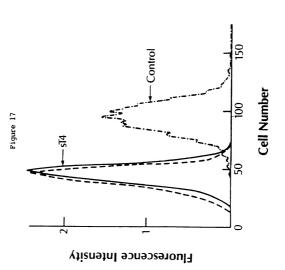


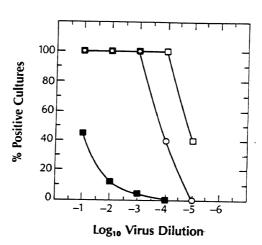
Figure 16

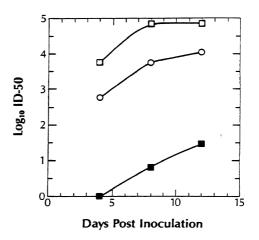


T4 +370
— thr pro val gin pro met ala ieu —
— ACC CCG GTC CAG CCA ATG GCC CTG —

8T4







30/30 Figure 18C

